

**Amendments to the specification:**

*Rewrite page 15, paragraph 5, as:*

Figure 3(a) ~~SEQ ID NO.: 339737 Gen Bank Accession No. M10988~~ (SEQ ID NO: 43) illustrates the DNA sequence encoding tumor necrosis factor (TNF $\alpha$ ) having the amino acid sequence shown in Fig. 3(b).

*Rewrite page 16, paragraph 2, as:*

Figure 3(b) ~~SEQ ID NO.: 339737 Gen Bank Accession No. M10988~~ (SEQ ID NO: 44) shows the amino acid sequence of human TNF $\alpha$  including the -76--1 presequence.

*Rewrite the paragraph bridging pages 37 and 38 as:*

The amino acid sequence of the P2 epitope is QYIKANSKFIGITEL (residues 225 to 239 of SEQ ID NO: 3) and corresponds to TT amino acids 830-844, and the sequence of the P30 epitope is FNNFTVSFWLRVPKVSASHLE (residues 224 to 244 of SEQ ID NO: 15) and corresponds to TT amino acids 947-967. Substituting P2 and P30 into two different human TNF $\alpha$  molecules would exchange approximately 10 % and 15 %, respectively, of the native TNF $\alpha$  sequence. In case both epitopes were inserted into a single TNF $\alpha$  molecule, about 25 % of the molecule would be exchanged, and one could fear that this would interfere too much with the remaining native parts of the TNF $\alpha$  molecule. It was therefore decided to develop two TNF $\alpha$  molecules, each containing either

P2 or P30. Together, such two molecules would be expected to be immunogenic in at least 80 % of the human population. In addition, it is very likely that truncated molecules composed partly of the P2 or P30 epitope and partly of TNF $\alpha$  flanking regions also will contribute to the immunogenicity resulting in the constructs being immunogenic in almost 100 % of the population.